SEQUENCE LISTING

- <110> Messier, Walter Sikela, James M
- <120> Methods to Identify Polynucleotide and Polypeptide Sequences Which May Be Associated with Physiological and Medical Conditions
- <130> GENO 200.2/CIP
- <140>
- <141>
- <150> 09/591,435
- <151> 2000-06-09
- <150> 09/240,915
- <151> 1999-01-29
- <150> 60/073,263
- <151> 1998-01-30
- <150> 60/098,987
- <151> 1998-09-02
- <160> 30
- <170> PatentIn Ver. 2.0
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- Cys Tyr Ser Asn Cys Pro Asp Gly Gln Ser Thr Ala Lys Thr Phe Leu 65 70 75 80
- Thr Val Tyr Trp Thr Pro Glu Arg Val Glu Leu Ala Pro Leu Pro Ser 85 90 95
- Trp Gln Pro Val Gly Lys Asp Leu Thr Leu Arg Cys Gln Val Glu Gly
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- Gly Ala Pro Arg Ala Asn Leu Thr Val Val Leu Leu Arg Gly Glu Lys
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- Thr Val Leu Val Glu Arg Asp His His Gly Ala Asn Phe Ser Cys Arg 145 150 155 160
- Thr Glu Leu Asp Leu Arg Pro Gln Gly Leu Gln Leu Phe Glu Asn Thr 165 170 175
- Ser Ala Pro His Gln Leu Gln Thr Phe Val Leu Pro Ala Thr Pro Pro 180 185 190
- Gln Leu Val Ser Pro Arg Val Leu Glu Val Asp Thr Gln Gly Thr Val 195 200 205
- Val Cys Ser Leu Asp Gly Leu Phe Pro Val Ser Glu Ala Gln Val His 210 215 220
- Leu Ala Leu Gly Asp Gln Arg Leu Asn Pro Thr Val Thr Tyr Gly Asn 225 230 235 240
- Asp Ser Phe Ser Ala Lys Ala Ser Val Ser Val Thr Ala Glu Asp Glu 245 250 255
- Gly Thr Gln Arg Leu Thr Cys Ala Val Ile Leu Gly Asn Gln Ser Arg 260 265 270
- Glu Thr Leu Gln Thr Val Thr Ile Tyr Ser Phe Pro Ala Pro Asn Val 275 280 285

- Ile Leu Thr Lys Pro Glu Val Ser Glu Gly Thr Glu Val Thr Val Lys 290 295 300
- Cys Glu Ala His Pro Arg Ala Lys Val Thr Leu Asn Gly Val Pro Ala 305 310 315 320
- Gln Pro Val Gly Pro Arg Val Gln Leu Leu Leu Lys Ala Thr Pro Glu 325 330 335
- Asp Asn Gly Arg Ser Phe Ser Cys Ser Ala Thr Leu Glu Val Ala Gly 340 345 350
- Gln Leu Ile His Lys Asn Gln Thr Arg Glu Leu Arg Val Leu Tyr Gly 355 360 365
- Pro Arg Leu Asp Glu Arg Asp Cys Pro Gly Asn Trp Thr Trp Pro Glu 370 375 380
- Asn Ser Gln Gln Thr Pro Met Cys Gln Ala Ser Gly Asn Pro Leu Pro 385 390 395 400
- Glu Leu Lys Cys Leu Lys Asp Gly Thr Phe Pro Leu Pro Val Gly Glu 405 410 415
- Ser Val Thr Val Thr Arg Asp Leu Glu Gly Thr Tyr Leu Cys Arg Ala 420 425 430
- Arg Ser Thr Gln Gly Glu Val Thr Arg Lys Val Thr Val Asn Val Leu 435 440 445
- Ser Pro Arg Tyr Glu Ile Val Ile Ile Thr Val Val Ala Ala Ala Val 450 455 460
- Ile Met Gly Thr Ala Gly Leu Ser Thr Tyr Leu Tyr Asn Arg Gln Arg 465 470 475 480
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<212> PRT

<213> Homo sapiens

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Arg Lys Val Tyr Glu Leu Ser Asn Val Gln Glu Asp Ser Gln Pro Met 50 55 60

Cys Tyr Ser Asn Cys Pro Asp Gly Gln Ser Thr Ala Lys Thr Phe Leu 65 70 75 80

Thr Val Tyr Trp Thr Pro Glu Arg Val Glu Leu Ala Pro Leu Pro Ser 85 90 95

Trp Gln Pro Val Gly Lys Asn Leu Thr Leu Arg Cys Gln Val Glu Gly
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Gly Ala Pro Arg Ala Asn Leu Thr Val Val Leu Leu Arg Gly Glu Lys
115 120 125

Glu Leu Lys Arg Glu Pro Ala Val Gly Glu Pro Ala Glu Val Thr Thr

130 135 140

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Ser	Ala	Pro	Tyr 180	Gln	Leu	Gln	Thr	Phe 185	Val	Leu	Pro	Ala	Thr 190	Pro	Pro
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Val	Cys 210	Ser	Leu	Asp	Gly	Leu 215	Phe	Pro	Val	Ser	Glu 220	Ala	Gln	Val	His
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Asp	Ser	Phe	Ser	Ala 245	Lys	Ala	Ser	Val	Ser 250	Val	Thr	Ala	Glu	Asp 255	Glu
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Glu	Thr	Leu 275		Thr	Val	Thr	Ile 280	Tyr	Ser	Phe	Pro	Ala 285	Pro	Asn	Val
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Cys		ı Ala	His		Arg				Thr	Leu 315		Gly	Val	Pro	Ala 320
Gln	n Pro	Leu	ı Gly	7 Pro		Ala	Gln	Leu	Leu 330		Lys	Ala	Thr	Pro 335	Glu
Asp) Asr	ı Gly	7 Arg		. Phe	e Ser	Cys	Ser 345		Thr	Leu	Glu	Val		Gly
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Glu Leu Lys Cys Leu Lys Asp Gly Thr Phe Pro Leu Pro Ile Gly Glu 410 405 415

Ser Val Thr Val Thr Arg Asp Leu Glu Gly Thr Tyr Leu Cys Arg Ala 420 425 430

Arg Ser Thr Gln Gly Glu Val Thr Arg Glu Val Thr Val Asn Val Leu 440 445 435

Ser Pro Arq Tyr Glu Ile Val Ile Ile Thr Val Val Ala Ala Ala Val 455 450

Ile Met Gly Thr Ala Gly Leu Ser Thr Tyr Leu Tyr Asn Arg Gln Arg 470 480 465

Lys Ile Lys Lys Tyr Arg Leu Gln Gln Ala Gln Lys Gly Thr Pro Met 485 490 495

Lys Pro Asn Thr Gln Ala Thr Pro Pro 505 500

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Gln Pro Glu Val Gly Gly Leu Glu Thr Ser Leu Asp Lys Ile Leu Leu 40

Asp Glu Gln Ala Gln Trp Lys His Tyr Leu Val Ser Asn Ile Ser His 55

Asp Thr Val Leu Gln Cys His Phe Thr Cys Ser Gly Lys Gln Glu Ser 75 70

Met Asn Ser Asn Val Ser Val Tyr Gln Pro Pro Arg Gln Val Ile Leu 90 85

Thr Leu Gln Pro Thr Leu Val Ala Val Gly Lys Ser Phe Thr Ile Glu 100 105 110

Cys Arg Val Pro Thr Val Glu Pro Leu Asp Ser Leu Thr Leu Phe Leu 115 120 125

Phe Arg Gly Asn Glu Thr Leu His Tyr Glu Thr Phe Gly Lys Ala Ala 130 135 140

Pro Ala Pro Gln Glu Ala Thr Ala Thr Phe Asn Ser Thr Ala Asp Arg 145 150 155 160

Glu Asp Gly His Arg Asn Phe Ser Cys Leu Ala Val Leu Asp Leu Met 165 170 175

Ser Arg Gly Gly Asn Ile Phe His Lys His Ser Ala Pro Lys Met Leu 180 185 190

Glu Ile Tyr Glu Pro Val Ser Asp Ser Gln Met Val Ile Ile Val Thr 195 200 205

Val Val Ser Val Leu Leu Ser Leu Phe Val Thr Ser Val Leu Leu Cys 210 215 220

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<212> PRT

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Met Gly Trp Ala Ala Phe Asn Leu Ser Asn Val Thr Gly Asn Ser Arg 50 55 60

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- Glu Gly Gly Ser Pro Arg Thr Ser Leu Thr Val Val Leu Leu Arg Trp 115 120 125
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- Thr Ala Thr Val Leu Ala Ser Arg Asp Asp His Gly Ala Pro Phe Ser 145 150 155 160
- Cys Arg Thr Glu Leu Asp Met Gln Pro Gln Gly Leu Gly Leu Phe Val 165 170 175
- Asn Thr Ser Ala Pro Arg Gln Leu Arg Thr Phe Val Leu Pro Val Thr 180 185 190
- Pro Pro Arg Leu Val Ala Pro Arg Phe Leu Glu Val Glu Thr Ser Trp 195 200 205
- Pro Val Asp Cys Thr Leu Asp Gly Leu Phe Pro Ala Ser Glu Ala Gln 210 215 220
- Val Tyr Leu Ala Leu Gly Asp Gln Met Leu Asn Ala Thr Val Met Asn 225 230 235 240
- His Gly Asp Thr Leu Thr Ala Thr Ala Thr Ala Thr Ala Arg Ala Asp 245 250 255
- Gln Glu Gly Ala Arg Glu Ile Val Cys Asn Val Thr Leu Gly Gly Glu 260 265 270
- Arg Arg Glu Ala Arg Glu Asn Leu Thr Val Phe Ser Phe Leu Gly Pro 275 280 285
- Ile Val Asn Leu Ser Glu Pro Thr Ala His Glu Gly Ser Thr Val Thr 290 295 300
- Val Ser Cys Met Ala Gly Ala Arg Val Gln Val Thr Leu Asp Gly Val 305 310 315 320

Pro Ala Ala Ala Pro Gly Gln Pro Ala Gln Leu Gln Leu Asn Ala Thr 325 330 335

Glu Ser Asp Asp Gly Arg Ser Phe Phe Cys Ser Ala Thr Leu Glu Val

Asp Gly Glu Phe Leu His Arg Asn Ser Ser Val Gln Leu Arg Val Leu 355 360 365

Tyr Gly Pro Lys Ile Asp Arg Ala Thr Cys Pro Gln His Leu Lys Trp 370 375 380

Lys Asp Lys Thr Arg His Val Leu Gln Cys Gln Ala Arg Gly Asn Pro 385 390 395 400

Tyr Pro Glu Leu Arg Cys Leu Lys Glu Gly Ser Ser Arg Glu Val Pro 405 410 415

Val Gly Ile Pro Phe Phe Val Asn Val Thr His Asn Gly Thr Tyr Gln 420 425 430

Cys Gln Ala Ser Ser Ser Arg Gly Lys Tyr Thr Leu Val Val Met
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Asp Ile Glu Ala Gly Ser Ser His Phe Val Pro Val Phe Val Ala Val 450 455 460

Leu Leu Thr Leu Gly Val Val Thr Ile Val Leu Ala Leu Met Tyr Val 465 470 475 480

Phe Arg Glu His Gln Arg Ser Gly Ser Tyr His Val Arg Glu Glu Ser 485 490 495

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Gly Leu Gln Val Tyr Asn Lys Cys Trp Lys Leu Glu His Cys Asn Phe 35 40 45

Lys Asp Leu Thr Thr Arg Leu Arg Glu Asn Glu Leu Thr Tyr Tyr Cys 50 55 60

Cys Lys Lys Asp Leu Cys Asn Phe Asn Glu Gln Leu Glu Asn Gly Gly 65 70 75 80

Asn Glu Gln Leu Glu Asn Gly Gly Asn Glu Gln Leu Glu Asn Gly Gly 85 90 95

Thr Ser Leu Ser Glu Lys Thr Val Leu Leu Arg Val Thr Pro Phe Leu 100 105 110

Ala Ala Ala Trp Ser Leu His Pro 115 120

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<211> 5140

<212> DNA

<213> Homo sapiens

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<213> Homo sapiens

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105

90

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85

100

95

110

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			_		_					ctc Leu			850
_										acc Thr			898
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	_									ctg Leu			994
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	_		_		-					ggc Gly 255			1186
				_						gtg Val			1234
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	Pro								cgg Arg 990					3394
Glu									Glu				tgc Cys 1010	3442
				Pro				Glu					gcg Ala	3490
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	_	_	_											ggc		3730
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_	_	_	_	_	_	_								gcc		3874
	Met L140	1111	Ala	пур		1145	PHE	AIA	MEC		цец 1150	Asp	FIO	Ala	AIA	
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	_													Arg		
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Thr	Val	Ser			Pro	Thr	Ser			Ser	Ile	Thr	His	Val	Ser	
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vab	Ser		1190	Giu	ber	БуЗ		1195	110	Oru	ALG		1200		017	
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Glu	Ser	Lys	Glu	Ala												
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<213> Homo sapiens

<400> 16

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35 40 45

Ser Cys Arg Val Ala Glu Ser Met Ala Pro Asp Pro Arg Thr Leu Gln 50 55 60

Arg Met Ala Cys Glu Val Ala Cys Gly Val Leu His Leu His Arg Asn 65 70 75 80

Asn Phe Val His Ser Asp Leu Ala Leu Arg Asn Cys Leu Leu Thr Ala 85 90 95 Asp Leu Thr Val Lys Ile Gly Asp Tyr Gly Leu Ala His Cys Lys Tyr Arg Glu Asp Tyr Phe Val Thr Ala Asp Gln Leu Trp Val Pro Leu Arg Trp Ile Ala Pro Glu Leu Val Asp Glu Val His Ser Asn Leu Leu Val Val Asp Gln Thr Lys Ser Gly Asn Val Trp Ser Leu Gly Val Thr Ile Trp Glu Leu Phe Glu Leu Gly Thr Gln Pro Tyr Pro Gln His Ser Asp Gln Gln Val Leu Ala Tyr Thr Val Arg Glu Gln Gln Leu Lys Leu Pro Lys Pro Gln Leu Gln Leu Thr Leu Ser Asp Arg Trp Tyr Glu Val Met Gln Phe Cys Trp Leu Gln Pro Glu Gln Arg Pro Thr Ala Glu Glu Val His Leu Leu Ser Tyr Leu Cys Ala Lys Gly Ala Thr Glu Ala Glu Glu Glu Phe Glu Arg Arg Trp Arg Ser Leu Arg Pro Gly Gly Gly Gly Val Gly Pro Gly Pro Gly Ala Ala Gly Pro Met Leu Gly Gly Val Val Glu Leu Ala Ala Ser Ser Phe Pro Leu Leu Glu Gln Phe Ala Gly Asp Gly Phe His Ala Asp Gly Asp Asp Val Leu Thr Val Thr Glu Thr Ser Arg Gly Leu Asn Phe Glu Tyr Lys Trp Glu Ala Gly Arg Gly Ala Glu Ala Phe Pro Ala Thr Leu Ser Pro Gly Arg Thr Ala Arg Leu Gln 

Glu Leu Cys Ala Pro Asp Gly Ala Pro Pro Gly Val Val Pro Val Leu

- Ser Ala His Ser Pro Ser Leu Gly Ser Glu Tyr Phe Ile Arg Leu Glu 355 360 365
- Glu Ala Ala Pro Ala Ala Gly His Asp Pro Asp Cys Ala Gly Cys Ala 370 375 380
- Pro Ser Pro Pro Ala Thr Ala Asp Gln Asp Asp Asp Ser Asp Gly Ser 385 390 395 400
- Thr Ala Ala Ser Leu Ala Met Glu Pro Leu Leu Gly His Gly Pro Pro 405 410 415
- Val Asp Val Pro Trp Gly Arg Gly Asp His Tyr Pro Arg Arg Ser Leu 420 425 430
- Ala Arg Asp Pro Leu Cys Pro Ser Arg Ser Pro Ser Pro Ser Ala Gly
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- Pro Leu Ser Leu Ala Glu Gly Gly Ala Glu Asp Ala Asp Trp Gly Val 450 455 460
- Ala Ala Phe Cys Pro Ala Phe Phe Glu Asp Pro Leu Gly Thr Ser Pro 465 470 475 480
- Leu Gly Ser Ser Gly Ala Pro Pro Leu Pro Leu Thr Gly Glu Asp Glu
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- Leu Glu Glu Val Gly Ala Arg Arg Ala Ala Gln Arg Gly His Trp Arg 500 505 510
- Ser Asn Val Ser Ala Asn Asn Ser Gly Ser Arg Cys Pro Glu Ser 515 520 525
- Trp Asp Pro Val Ser Ala Gly Cys His Ala Glu Gly Cys Pro Ser Pro 530 535 540
- Lys Gln Thr Pro Arg Ala Ser Pro Glu Pro Gly Tyr Pro Gly Glu Pro 545 550 555 560
- Leu Leu Gly Leu Gln Ala Ala Ser Ala Gln Glu Pro Gly Cys Cys Pro
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- Gly Leu Pro His Leu Cys Ser Ala Gln Gly Leu Ala Pro Ala Pro Cys 580 585 590
- Leu Val Thr Pro Ser Trp Thr Glu Thr Ala Ser Ser Gly Gly Asp His 595 600 605

Pro Gln Ala Glu Pro Lys Leu Ala Thr Glu Ala Glu Gly Thr Thr Gly Pro Arg Leu Pro Leu Pro Ser Val Pro Ser Pro Ser Gln Glu Gly Ala Pro Leu Pro Ser Glu Glu Ala Ser Ala Pro Asp Ala Pro Asp Ala Leu Pro Asp Ser Pro Thr Pro Ala Thr Gly Gly Glu Val Ser Ala Ile Lys Leu Ala Ser Ala Leu Asn Gly Ser Ser Ser Pro Glu Val Glu Ala Pro Ser Ser Glu Asp Glu Asp Thr Ala Glu Ala Thr Ser Gly Ile Phe Thr Asp Thr Ser Ser Asp Gly Leu Gln Ala Arg Arg Pro Asp Val Val Pro Ala Phe Arg Ser Leu Gln Lys Gln Val Gly Thr Pro Asp Ser Leu Asp Ser Leu Asp Ile Pro Ser Ser Ala Ser Asp Gly Gly Tyr Glu Val Phe Ser Pro Ser Ala Thr Gly Pro Ser Gly Gly Gln Pro Arg Ala Leu Asp Ser Gly Tyr Asp Thr Glu Asn Tyr Glu Ser Pro Glu Phe Val Leu Lys Glu Ala Gln Glu Gly Cys Glu Pro Gln Ala Phe Ala Glu Leu Ala Ser Glu Gly Glu Gly Pro Gly Pro Glu Thr Arg Leu Ser Thr Ser Leu Ser Gly Leu Asn Glu Lys Asn Pro Tyr Arg Asp Ser Ala Tyr Phe Ser Asp Leu Glu Ala Glu Ala Glu Ala Thr Ser Gly Pro Glu Lys Lys Cys Gly Gly Asp Arg Ala Pro Gly Pro Glu Leu Gly Leu Pro Ser Thr Gly

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Gln	Gly	Ser	Gly	Pro 885	Gly	Glu	Val	Leu	Pro 890	Pro	Leu	Leu	Gln	Leu 895	Glu
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Pro	Pro	Glu 915	Pro	Gln	Gly	Pro	Ala 920	Lys	Val	Arg	Pro	Gly 925	Pro	Ser	Pro
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Arg	Leu	Ala	Leu 980	Pro	Gly	Leu	Pro	Ala 985	Ala	Leu	Glu	Gly	Arg 990	Pro	Glu
Glu	Glu	Glu 995	Glu	Asp	Ser		Asp 1000	Ser	Asp	Glu		Asp 1005	Glu	Glu	Leu
_	Cys 1010	_	Ser	Val		Glu 1015	Pro	Ser	Glu		Ser 1020	Glu	Glu	Glu	Ala
Pro 025	Ala	Val	Pro		Val 1030	Val	Ala	Glu		Gln 1035	Ser	Ala	Arg		Leu 1040
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Phe Pro Leu Met Thr Ala Lys Ala Ala Phe Ala Met Ala Leu Asp Pro 1140 1145 1150

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